

[illegible]

1

<211> 9  
<212> PRT  
<213> Histocompatibility antigen

<220>  
<221> PEPTIDE  
<222> (2)  
<223> AMINO ACID X REPRESENTS EITHER ISOLEUCINE OR  
LEUCINE RESIDUES

<400> 3  
Tyr Xaa Thr Asp Arg Val Met Thr Val  
1 5

<210> 4  
<211> 9  
<212> PRT  
<213> Histocompatibility antigen

<220>  
<221> PEPTIDE  
<222> (2)..(7)  
<223> AMINO ACIDS X REPRESENT LEUCINE OR ISOLEUCINE  
RESIDUES

<400> 4  
Val Xaa His Asp Asp Xaa Xaa Glu Ala  
1 5

Sub A

<210> 5  
<211> 9  
<212> PRT  
<213> KIAA0223 partial complementary DNA

<400> 5  
Val Leu Arg Asp Asp Leu Leu Glu Ala  
1 5

<210> 6  
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<212> DNA  
<213> Unknown Organism

<220>  
<223> Description of Unknown Organism:Reverse primer

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25

<210> 7  
<211> 24  
<212> DNA  
<213> Unknown Organism

<220>  
<223> Description of Unknown Organism:Forward primer

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24

<210> 8  
<211> 25  
<212> DNA  
<213> Unknown Organism

<220>  
<223> Description of Unknown Organism:Reverse Primer

<400> 8  
gaaggccaca gcaatcgtct ccagg

25

<210> 9  
<211> 30  
<212> DNA  
<213> Unknown Organism

<220>  
<223> Description of Unknown Organism:HA-1" allele  
primer H1

<400> 9  
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30

<210> 10  
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<212> DNA  
<213> Unknown Organism

<220>

*Sub A*

<223> Description of Unknown Organism:HA-1r allele  
primer R1

<400> 10

ccttgagaaa cttaaggagt gtgtgttgcg

30

<210> 11

<211> 33

<212> DNA

<213> Forward KIAA00223 based PCR primer

<400> 11

ccggcatgga cgtcgtcgag gacatctccc atc

33

<210> 12

<211> 30

<212> DNA

<213> Reverse KIAA00223 based PCR primer

<400> 12

ctacttcagg ccacagcaat cgtctccagg

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<210> 13

<211> 38

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism:HA-1-/-

<400> 13

gagtgtgtgt tgcgtgacga cctccttgag gcccgccg

38

<210> 14

<211> 13

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism:HA-1-/-

<400> 14

Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg

1

5

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Sub A

<210> 15  
<211> 38  
<212> DNA  
<213> Unknown Organism

<220>  
<223> Description of Unknown Organism:HA-1+/+

<400> 15  
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38

<210> 16  
<211> 13  
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<213> Unknown Organism

Sub A

<220>  
<223> Description of Unknown Organism:HA-1+/+

<400> 16  
Glu Cys Val Leu His Asp Asp Leu Leu Glu Ala Arg Arg  
1 5 10